SEQUENCE LISTING

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<120> Methods for Analysing Animal Products

<130> 62781.000013

<140> 10/758,422
<141> 2004-01-16

<150> 09/450,651
<151> 1999-11-30

<150> PCT/GB98/01531
<151> 1998-05-27

<150> GB 9801990
<151> 1998-01-31

<150> GB 9711214.8
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Arg Thr Glu Thr Pro Val
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Pro Met Tyr Tyr Phe Val Cys Cys Leu Ala Val Ser Asp Leu Leu Val
        35                  40                  45

Ser Val Ser Asn Val Leu Glu Thr Ala Val Leu Leu Leu Leu Glu Ala
    50                  55                  60

Gly Ala Leu Ala Ala Gln Ala Ala Val Val Gln Gln Leu Asp Asn Val
65                  70                  75                  80

Met Asp Val Leu Ile Cys Gly Ser Met Val Ser Ser Leu Cys Phe Leu
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Gly Ala Ile Ala Val Asp Arg Tyr Val Ser Ile Phe Tyr Ala Leu Arg
            100                 105                 110

Tyr His Ser Ile Val Thr Leu Pro Arg Ala Gly Arg Ala Ile Ala Ala
        115                 120                 125

Ile Trp Ala Gly Ser Val Leu Ser Ser Thr Leu Phe Ile Ala Tyr Tyr
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His His Thr Ala Val Leu Leu Gly Leu Val Ser Phe Phe Val Ala Met
145                 150                 155                 160

Leu Ala Leu Met Ala Val Leu Tyr Val His Met Leu Ala Arg Ala Cys
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Gln His Gly Arg His Ile Ala Arg Leu His Lys Thr Gln His Pro Thr
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Arg Gln Gly Cys Gly Leu Lys Gly Ala Ala Thr Leu Thr Ile Leu Leu
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Gly Val Phe Leu Leu Cys Trp Ala Pro Phe Phe Leu His Leu Ser Leu
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Val Val Leu Cys Pro Gln His Pro Thr Cys Gly Cys Val Phe Lys Asn
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225 230 235 240

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Asn Val Leu Val Val Ala Ala Ile Ala Lys Asn Arg Asn Leu His Ser
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Pro Met Tyr Tyr Phe Val Cys Cys Leu Ala Val Ser Asp Leu Leu Val
35 40 45

Ser Val Ser Asn Met Leu Glu Thr Ala Val Leu Pro Leu Leu Glu Ala
50 55 60

Gly Ala Leu Ala Ala Gln Ala Ala Val Val Gln Gln Leu Asp Asn Val
65 70 75 80

Met Asp Val Leu Ile Cys Gly Ser Met Val Ser Ser Leu Cys Phe Leu
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Gly Ala Ile Ala Val Asp Arg Tyr Val Ser Ile Phe Tyr Ala Leu Arg
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115 120 125

Ile Trp Ala Gly Ser Val Leu Ser Ser Thr Leu Phe Ile Ala Tyr Tyr
130 135 140

His His Thr Ala Val Leu Leu Gly Leu Val Ser Phe Phe Val Ala Met

145 150 155 160

Leu Ala Leu Met Ala Val Leu Tyr Val His Met Leu Ala Arg Ala Cys
165 170 175

Gln His Gly Arg His Ile Ala Arg Leu His Lys Thr Gln His Pro Thr
180 185 190

Arg Gln Gly Cys Gly Leu Lys Gly Ala Ala Thr Leu Thr Ile Leu Leu
195 200 205

Gly Val Phe Leu Leu Cys Trp Ala Pro Phe Phe Leu His Leu Ser Leu
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Val Val Leu Cys Pro Gln His Pro Thr Cys Gly Cys Val Phe Lys Asn
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Pro Met Tyr Tyr Phe Val Cys Cys Leu Ala Val Ser Asp Leu Leu Val
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Gly Ala Leu Ala Ala Gln Ala Ala Val Val Gln Gln Leu Asp Asn Val

65 70 75 80

Met Asn Val Leu Ile Cys Gly Ser Met Val Ser Ser Leu Cys Phe Leu
85 90 95

Gly Ala Ile Ala Val Asp Arg Tyr Val Ser Ile Phe Tyr Ala Leu Arg
100 105 110

Tyr His Ser Ile Val Thr Leu Pro Arg Ala Gly Arg Ala Ile Ala Ala
115 120 125

Ile Trp Ala Gly Ser Val Leu Ser Ser Thr Leu Phe Ile Ala Tyr Tyr
130 135 140

His His Thr Ala Val Leu Leu Gly Leu Val Ser Phe Phe Val Ala Met
145 150 155 160

Leu Ala Leu Met Ala Val Leu Tyr Val His Met Leu Ala Arg Ala Cys
165 170 175

Gln His Gly Arg His Ile Ala Arg Leu His Lys Thr Gln His Pro Thr
180 185 190

Arg Gln Gly Cys Gly Leu Lys Gly Ala Ala Thr Leu Thr Ile Leu Leu
195 200 205

Gly Val Phe Leu Leu Cys Trp Ala Pro Phe Phe Leu His Leu Ser Leu
210 215 220

Val Val Leu Cys Pro Gln His Pro Thr Cys Gly Cys Val Phe Lys Asn
225 230 235 240

Val Asn Leu Phe Leu Ala Leu Val Ile Cys Asn Ser Ile
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<210> 61
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<213> Sus sp.

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<400> 61

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Asn Val Leu Val Val Ala Ala Ile Ala Lys Asn Arg Asn Leu His Ser
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Pro Met Tyr Tyr Phe Val Cys Cys Leu Ala Val Ser Asp Leu Leu Val
35 40 45

Ser Val Ser Asn Val Leu Glu Thr Ala Val Leu Leu Leu Leu Glu Ala
50 55 60

Gly Ala Leu Ala Ala Gln Ala Ala Val Val Gln Gln Leu Asp Asn Val
65 70 75 80

Met Asp Val Leu Ile Cys Gly Ser Met Val Ser Ser Leu Cys Phe Leu
85 90 95

Gly Ala Ile Ala Val Asp Arg Tyr Val Ser Ile Phe Tyr Ala Leu Arg
100 105 110

Tyr His Ser Ile Val Thr Leu Pro Arg Val Gly Arg Ala Ile Ala Ala
115 120 125

Ile Trp Ala Gly Ser Val Leu Ser Ser Thr Leu Phe Ile Ala Tyr Tyr
130 135 140

His His Thr Ala Val Leu Leu Gly Leu Val Ser Phe Phe Val Ala Met
145 150 155 160

Leu Ala Leu Met Ala Val Leu Tyr Val His Met Leu Ala Arg Ala Cys
165 170 175

Gln His Gly Arg His Ile Ala Arg Leu His Lys Thr Gln His Pro Thr
180 185 190

Arg Gln Gly Cys Gly Leu Lys Gly Thr Ala Thr Leu Thr Ile Leu Leu
195 200 205

Gly Val Phe Leu Leu Cys Trp Ala Pro Phe Phe Leu His Leu Ser Leu
210 215 220

Val Val Leu Cys Pro Gln His Pro Thr Cys Gly Cys Val Phe Lys Asn
225 230 235 240

Val Asn Leu Phe Leu Ala Leu Val Ile Cys Asn Ser Ile
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ccagcaaaat cagagttaat cgtcagtgct ggcgatgaga ttaggctgtt ctgcaccgat 180

ccaggatctg tcaaatggac ttttgagacc ctgggtcagc tgagtgagaa tacacacgca 240

gagtggatcg tggagaaagc agaggccatg aatacaggca attatacatg caccaatgaa 300

ggcggtttaa gcagttccat ttatgtgttt gttagagatc ctgagaagct tttcctcgtc 360

gaccctccct tgtatgggaa ggaggacaat gacgcgctgg tccgatgtcc tctgacggac 420

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taccttctca gggaagggga ggaatttgcc gtgatgtgct tgatcaaaga cgtgtctagt 720

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gcacttccag gaggcatggg gagtgggggа ggctggagag acggcgggga gcgccacctc 240

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ggtgggacgt gtgcgggcag aagctgtccg ggtgttgctc actgcaggac ctcaggggaa 360

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gtgggacgtg tgcgggcaga agctgtccgg gtgttgctca ctgcaggacc tcaggggaag 360

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catccagaga ccaccacgcc cgcctttggg gcgcgctctg gggactttgc cccccactgg 300

ggtgggacgt gtgcgggcag aagctgtccg ggtgttgctc actgcaggac ctcaggggaa 360

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<210> 69
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<212> DNA
<213> Artificial Sequence

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